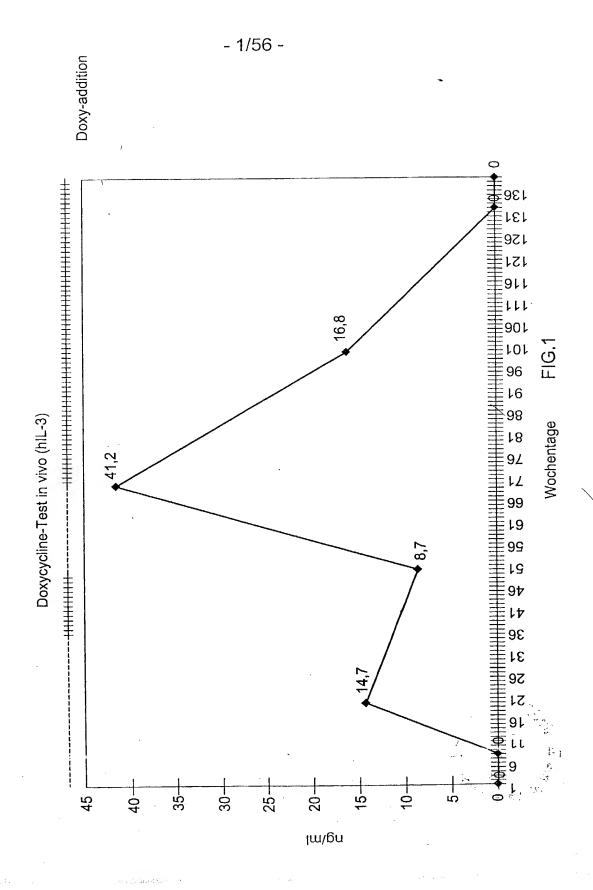
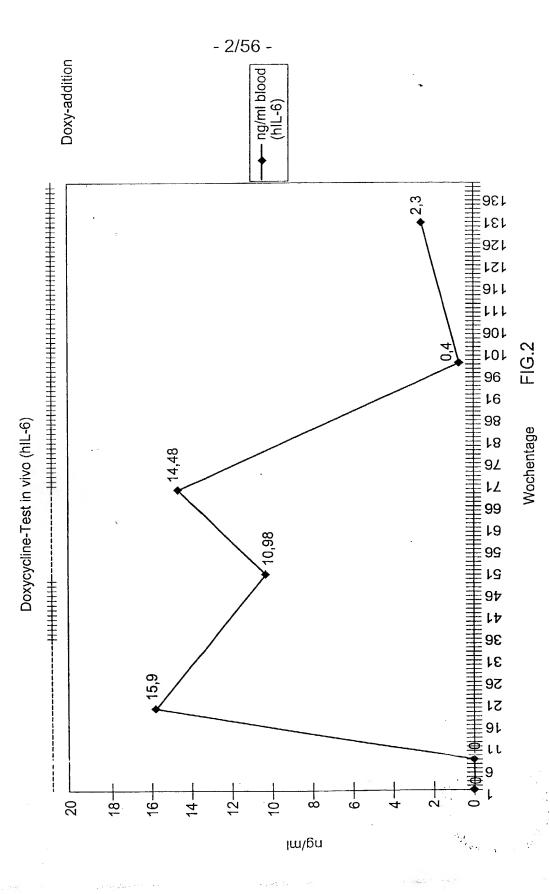
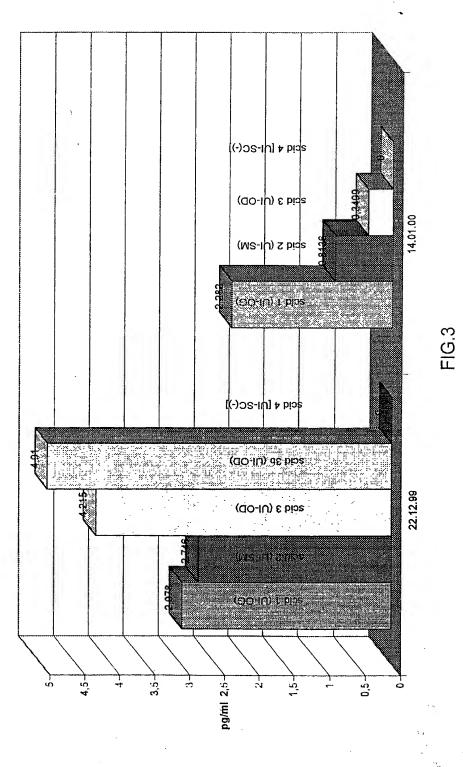


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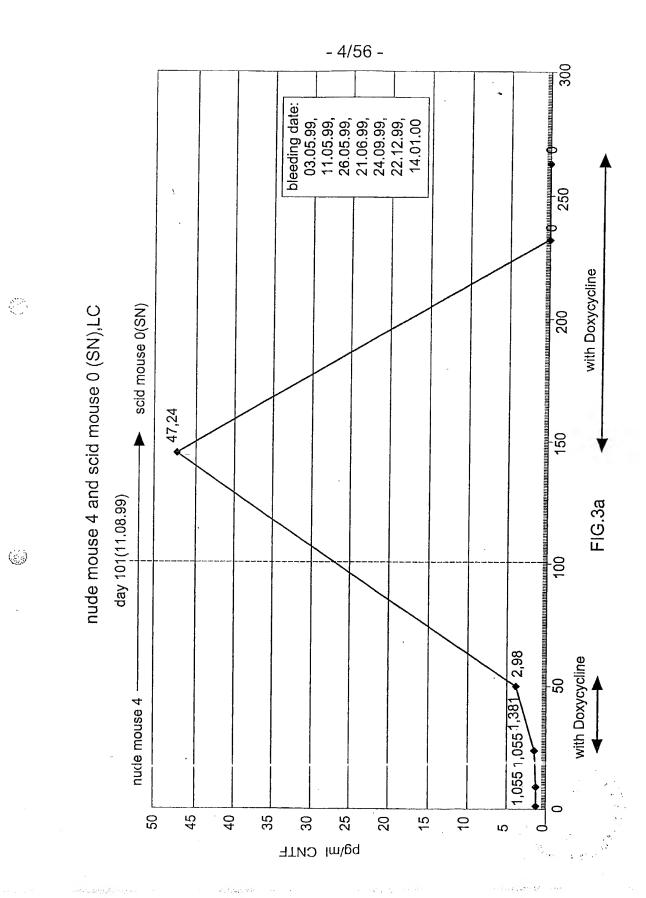




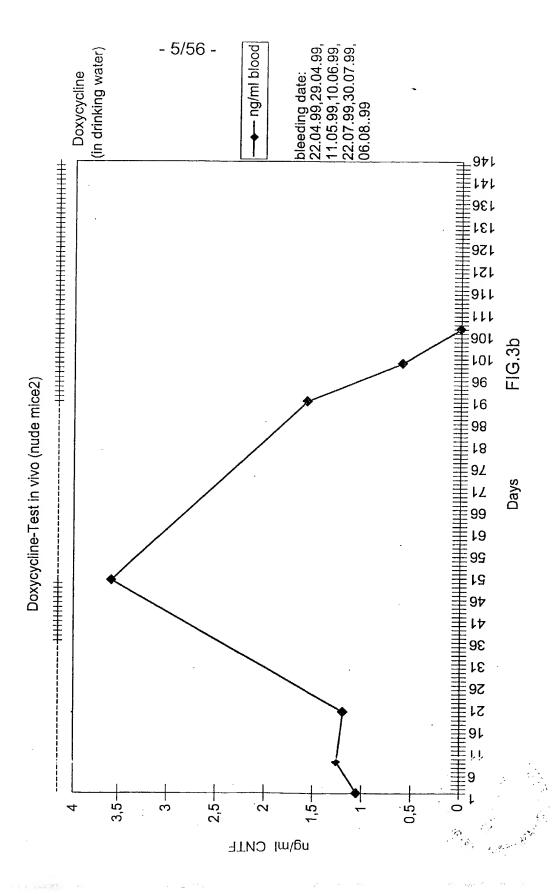
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Scid-mice [OG,SM,OD,SC(-)]:hIL-6



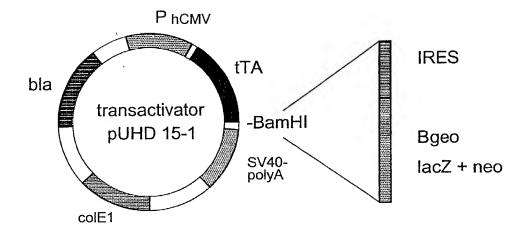
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- 6/56 - Cloning of growth factor genes



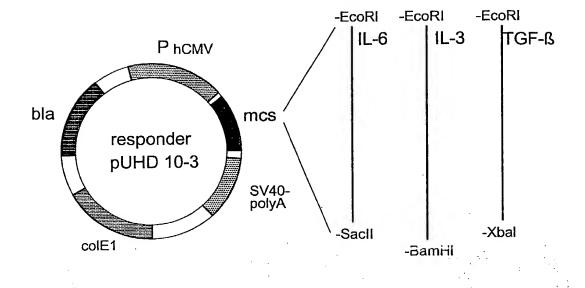
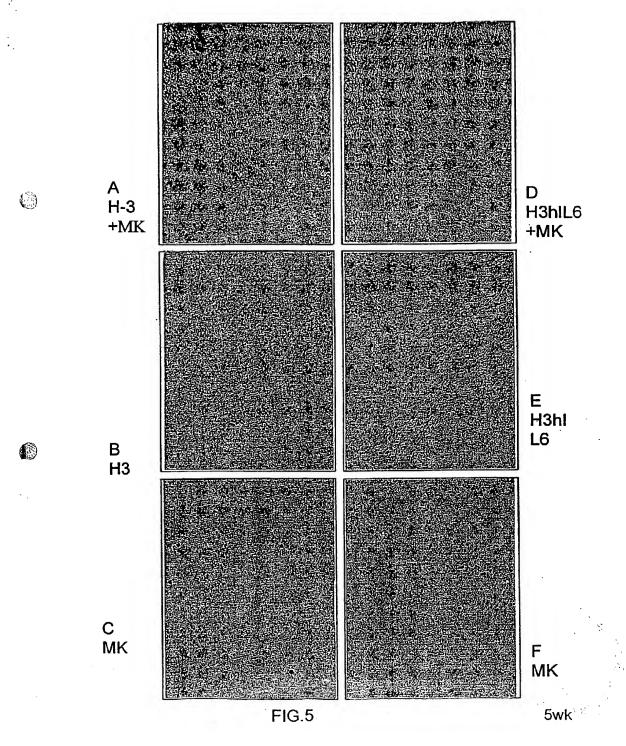
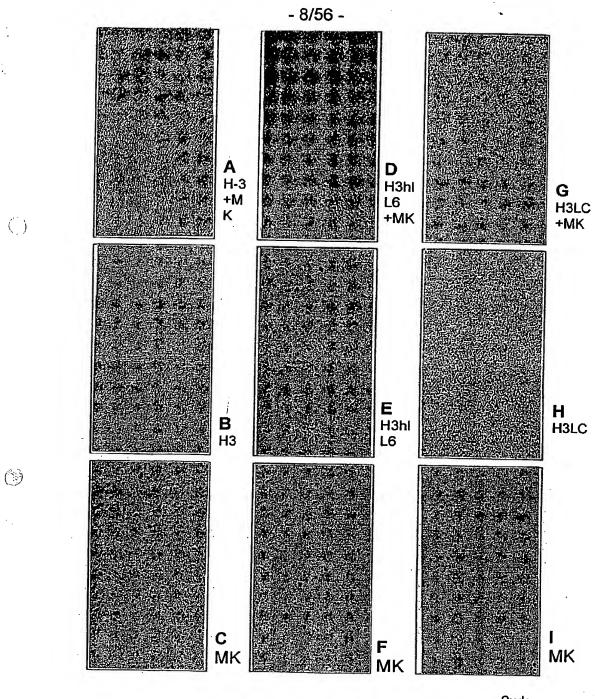


FIG.4

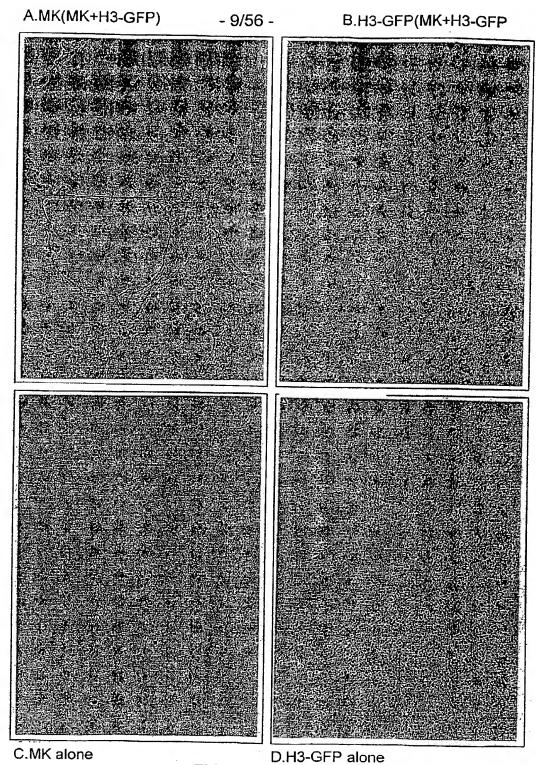






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FIG.6



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FIG.7

- 10/56 -A.MK (MK+H3-GFP-hIL6) B.H3-GFP-hIL6(MK+H3-GFP-hIL6)

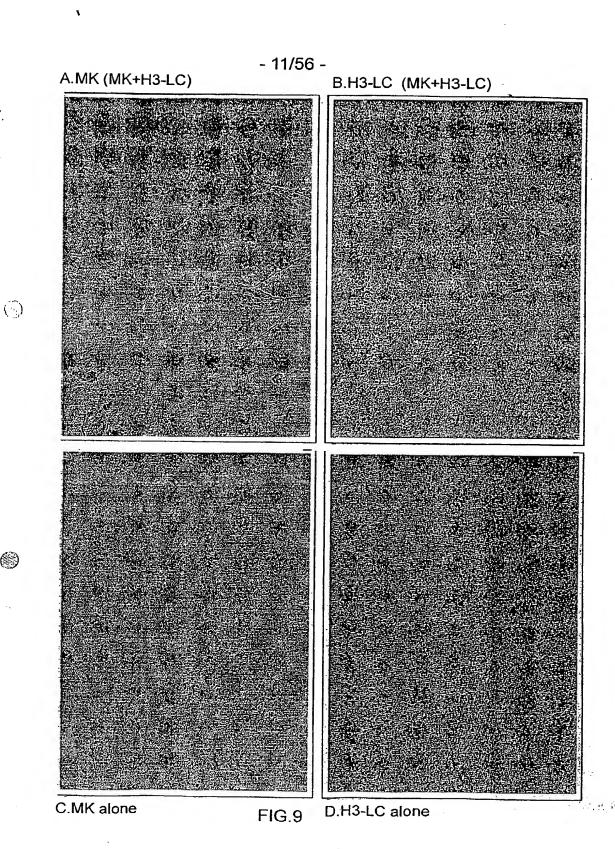
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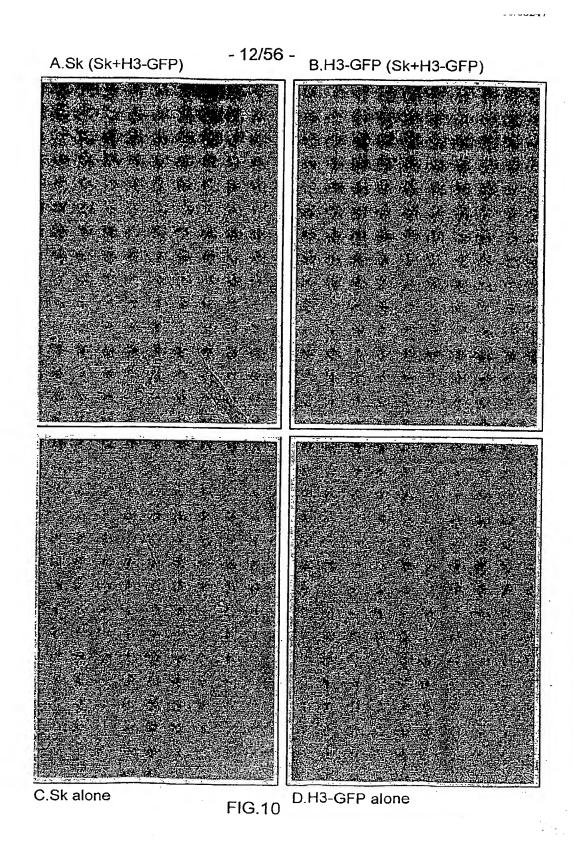
(3)

FIG.8

D.H3-GFP-hIL6 alone

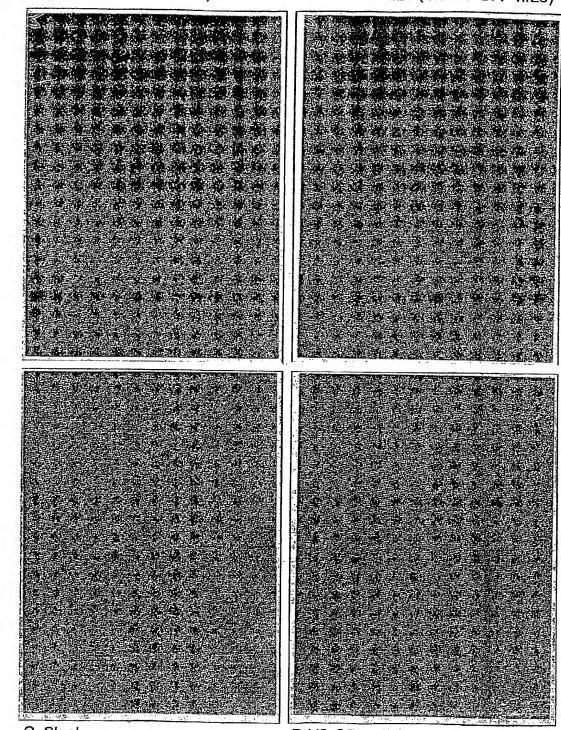
C.MK alone





()

# A. SK (Sk+H3-GFP-hIL6) - 13/56 - B. H3-GFP-hIL6 (Sk+H3-GFP-hIL6)



C. Sk alone

 $(\cdot)$ 

FIG.11

D.H3-GFP-hIL6 alone

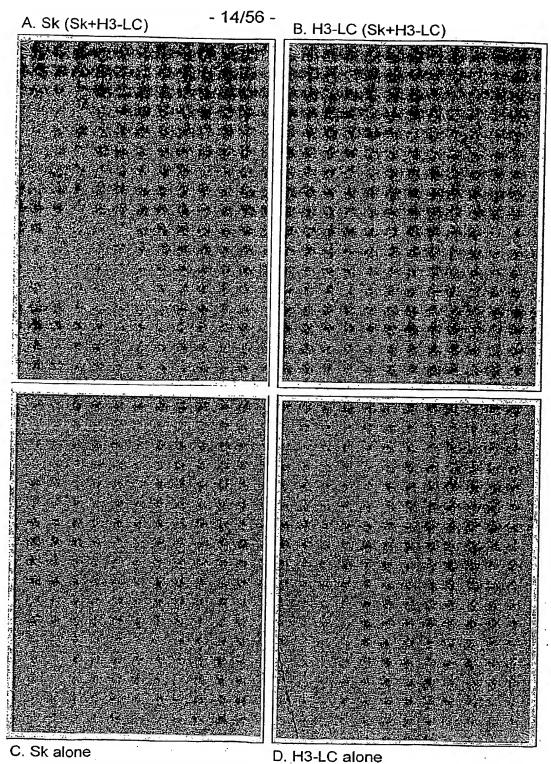
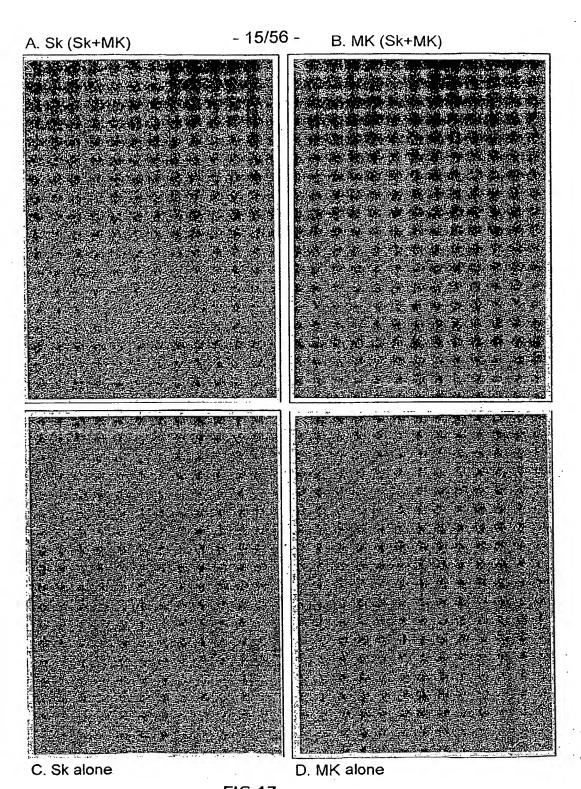
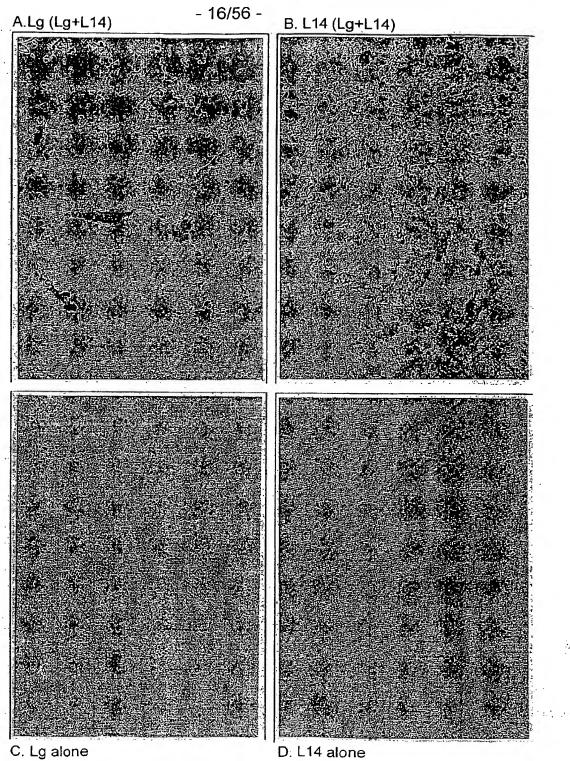


FIG.12

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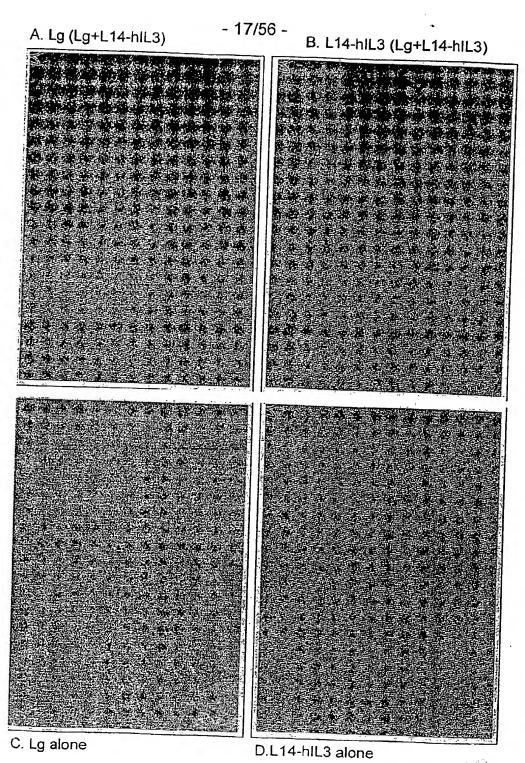
**FIG.13** 



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FIG.14



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(8)

FIG.15

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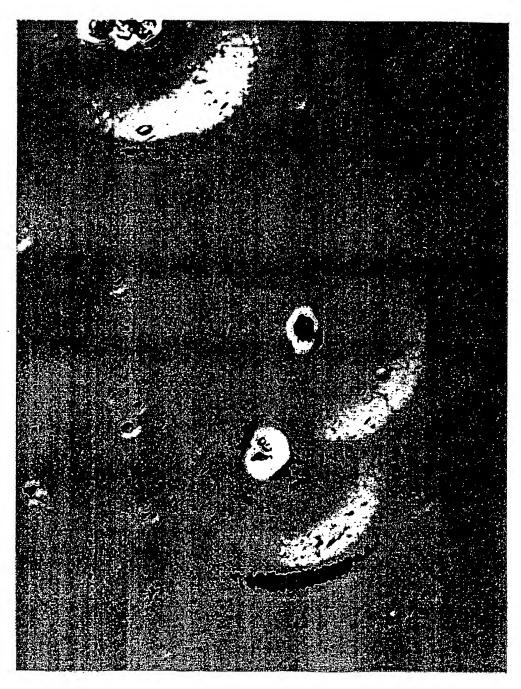


FIG.16

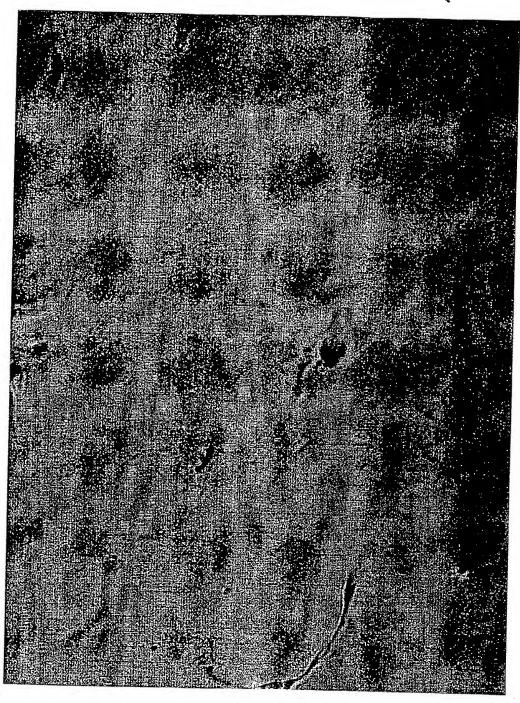


Fig. 17.

# - 20/56 -

# pD12JCVPlong-hCNTF

(i)

Length: 7969 July 22, 1999

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651	TGGGTTNTTC ATAAGCCATT GTTCTGAANT TTTTTAGCTT TGTAAATGAA
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1201	catcaacctg gactctgcgg atgggatgcc agtggcaagc actgatcagt
1251	ggagtgaget gaccgaggea gagcgactec aagagaacet teaagettat

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(45)

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### - 23/56 -

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5501 TCACGCTGTA GGTATCTCAG TTCGGTGTAG GTCGTTCGCT CCAAGCTGGG 5551 CTGTGTGCAC GAACCCCCCG TTCAGCCCGA CCGCTGCGCC TTATCCGGTA 5601 ACTATCGTCT TGAGTCCAAC CCGGTAAGAC ACGACTTATC GCCACTGGCA 5651 GCAGCCACTG GTAACAGGAT TAGCAGAGCG AGGTATGTAG GCGGTGCTAC 5701 AGAGTTCTTG AAGTGGTGGC CTAACTACGG CTACACTAGA AGGACAGTAT TTGGTATCTG CGCTCTGCTG AAGCCAGTTA CCTTCGGAAA AAGAGTTGGT AGCTCTTGAT CCGGCAAACA AACCACCGCT GGTAGCGGTG GTTTTTTTGT TTGCAAGCAG CAGATTACGC GCAGAAAAAA AGGATCTCAA GAAGATCCTT 5901 TGATCTTTC TACGGGGTCT GACGCTCAGT GGAACGAAAA CTCACGTTAA 5951 GGGATTTTGG TCATGAGATT ATCAAAAAGG ATCTTCACCT AGATCCTTTT 6001 AAATTAAAAA TGAAGTTTTA AATCAATCTA AAGTATATAT GAGTAAACTT 6051 GGTCTGACAG TTACCAATGC TTAATCAGTG AGGCACCTAT CTCAGCGATC 6101 TGTCTATTTC GTTCATCCAT AGTTGCCTGA CTCCCCGTCG TGTAGATAAC 6151 TACGATACGG GAGGGCTTAC CATCTGGCCC CAGTGCTGCA ATGATACCGC 6201 GAGACCCACG CTCACCGGCT CCAGATTAT CAGCAATAAA CCAGCCAGCC 6251 GGAAGGGCCG AGCGCAGAAG TGGTCCTGCA ACTTTATCCG CCTCCATCCA 6301 GTCTATTAAT TGTTGCCGGG AAGCTAGAGT AAGTAGTTCG CCAGTTAATA 6351 GTTTGCGCAA CGTTGTTGCC ATTGCTACAG GCATCGTGGT GTCACGCTCG TCGTTTGGTA TGGCTTCATT CAGCTCCGGT TCCCAACGAT CAAGGCGAGT TACATGATCC CCCATGTTGT GCAAAAAAGC GGTTAGCTCC TTCGGTCCTC CGATCGTTGT CAGAAGTAAG TTGGCCGCAG TGTTATCACT CATGGTTATG 6501 GCAGCACTGC ATAATTCTCT TACTGTCATG CCATCCGTAA GATGCTTTTC TGTGACTGGT GAGTACTCAA CCAAGTCATT CTGAGAATAG TGTATGCGGC 6601 GACCGAGTTG CTCTTGCCCG GCGTCAATAC GGGATAATAC CGCGCCACAT 6651 AGCAGAACTT TAAAAGTGCT CATCATTGGA AAACGTTCTT CGGGGCGAAA 6701 ACTCTCAAGG ATCTTACCGC TGTTGAGATC CAGTTCGATG TAACCCACTC GTGCACCCAA CTGATCTTCA GCATCTTTTA CTTTCACCAG CGTTTCTGGG 6851 TGAGCAAAAA CAGGAAGGCA AAATGCCGCA AAAAAGGGAA TAAGGGCGAC

### - 25/56 -

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Fig. 18

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     201
     251
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     301
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     401
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TCTAGAGGAT CTTTGTGAAG GAACCTTACT TCTGTGGTGT GACATAATTG
    2951
    3001
    3051
          GACAAACTAC CTACAGAGAT TTAAAGCTCT AAGGTAAATA TAAAATTTTT
    3101
          AAGTGTATAA TGTGTTAAAC TACTGATTCT AATTGTTTGT GTATTTTAGA
          TTCCAACCTA TGGAACTGAT GAATGGGAGC AGTGGTGGAA TGCCTTTAAT
    3151
    3201
          GAGGAAAACC TGTTTTGCTC AGAAGAAATG CCATCTAGTG ATGATGAGGC
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3251 TACTGCTGAC TCTCAACATT CTACTCCTCC AAAAAAGAAG AGAAAGGTAG AAGACCCCAA GGACTTTCCT TCAGAATTGC TAAGTTTTTT GAGTCATGCT GTGTTTAGTA ATAGAACTCT TGCTTGCTTT GCTATTTACA CCACAAAGGA 3351 AAAAGCTGCA CTGCTATACA AGAAAATTAT GGAAAAATAT TCTGTAACCT 3401 TTATAAGTAG GCATAACAGT TATAATCATA ACATACTGTT TTTTCTTACT CCACACAGGC ATAGAGTGTC TGCTATTAAT AACTATGCTC AAAAATTGTG 3451 3501 TACCTTTAGC TTTTTAATTT GTAAAGGGGT TAATAAGGAA TATTTGATGT 3551 3601 ATAGTGCCTT GACTAGAGAT CATAATCAGC CATACCACAT TTGTAGAGGT TTTACTTGCT TTAAAAAACC TCCCACACCT CCCCCTGAAC CTGAAACATA AAATGAATGC AATTGTTGTT GTTAACTTGT TTATTGCAGC TTATAATGGT 3651 3701 3751 TACAAATAAA GCAATAGCAT CACAAATTTC ACAAATAAAG CATTTTTTTC ACTGCATTCT AGTTGTGGTT TGTCCAAACT CATCAATGTA TCTTATCATG
TCTGGATCCC CGGGTCCCTA TAGTGAGTCG TATTAGCTTG GCGTAATCAT 3801 3851 3901 GGTCATAGCT GTTTCCTGTG TGAAATTGTT ATCCGCTCAC AATTCCACAC AACATACGAG CCGGAAGCAT AAAGTGTAAA GCCTGGGGTG CCTAATGAGT GAGCTAACTC ACATTAATTG CGTTGCGCTC ACTGCCCGCT TTCCAGTCGG 3951 4001 GAAACCTGTC GTGCCAGCTG CATTAATGAA TCGGCCAACG CGCGGGAGA GGCGGTTTGC GTATTGGGCG CTCTTCCGCT TCCTCGCTCA CTGACTCGCT GCGCTCGGTC GTTCGGCTGC GGCGAGCGGT ATCAGCTCAC TCAAAGGCGG 4051 4101 4151 4201 TAATACGGTT ATCCACAGAA TCAGGGGATA ACGCAGGAAA GAACATGTGA GCAAAAGGCC AGCAAAAGGC CAGGAACCGT AAAAAGGCCG CGTTGCTGGC GTTTTTCCAT AGGCTCCGCC CCCCTGACGA GCATCACAAA AATCGACGCT 4251 4301 4351 CAAGTCAGAG GTGGCGAAAC CCGACAGGAC TATAAAGATA CCAGGCGTTT CCCCCTGGAA GCTCCCTCGT GCGCTCTCCT GTTCCGACCC TGCCGCTTAC CGGATACCTG TCCGCCTTTC TCCCTTCGGG AAGCGTGGCG CTTTCTCAAT 4401 4451 4501 GCTCACGCTG TAGGTATCTC AGTTCGGTGT AGGTCGTTCG CTCCAAGCTG GGCTGTGTGC ACGAACCCCC CGTTCAGCCC GACCGCTGCG CCTTATCCGG TAACTATCGT CTTGAGTCCA ACCCGGTAAG ACACGACTTA TCGCCACTGG 4551 4601 CAGCAGCCAC TEGTAACAGG ATTAGCAGAG CGAGGTATGT AGGCGGTGCT
ACAGAGTTCT TGAAGTGGTG GCCTAACTAC GGCTACACTA GAAGGACAGT
ATTTGGTATC TGCGCTCTGC TGAAGCCAGT TACCTTCGGA AAAAGAGTTG 4651 4701 4751 4801 GTAGCTCTTG ATCCGGCAAA CAAACCACCG CTGGTAGCGG TGGTTTTTT GTTTGCAAGC AGCAGATTAC GCGCAGAAAA AAAGGATCTC AAGAAGATCC 4851 TTTGATCTTT TCTACGGGGT CTGACGCTCA GTGGAACGAA AACTCACGTT 4901 4951 AAGGGATTTT GGTCATGAGA TTATCAAAAA GGATCTTCAC CTAGATCCTT TTAAATTAAA AATGAAGTTT TAAATCAATC TAAAGTATAT ATGAGTAAAC TTGGTCTGAC AGTTACCAAT GCTTAATCAG TGAGGCACCT ATCTCAGCGA 5001 5051 5101 TCTGTCTATT TCGTTCATCC ATAGTTGCCT GACTCCCCGT CGTGTAGATA ACTACGATAC GGGAGGGCTT ACCATCTGGC CCCAGTGCTG CAATGATACC GCGAGACCCA CGCTCACCGG CTCCAGATTT ATCAGCAATA AACCAGCCAG 5151 5201 5251 CCGGAAGGGC CGAGCGCAGA AGTGGTCCTG CAACTTTATC CGCCTCCATC 5301 CAGTCTATTA ATTGTTGCCG GGAAGCTAGA GTAAGTAGTT CGCCAGTTAA 5351 TAGTTTGCGC AACGTTGTTG CCATTGCTAC AGGCATCGTG GTGTCACGCT 5401 CGTCGTTTGG TATGGCTTCA TTCAGCTCCG GTTCCCAACG ATCAAGGCGA GTTACATGAT CCCCCATGTT GTGCAAAAAA GCGGTTAGCT CCTTCGGTCC 5451 TCCGATCGTT GTCAGAAGTA AGTTGGCCGC AGTGTTATCA CTCATGGTTA 5501 TGGCAGCACT GCATAATTCT CTTACTGTCA TGCCATCCGT AAGATGCTTT
TCTGTGACTG GTGAGTACTC AACCAAGTCA TTCTGAGAAT AGTGTATGCG 5551 5601 5651 GCGACCGAGT TGCTCTTGCC CGGCGTCAAT ACGGGATAAT ACCGCGCCAC ATAGCAGAAC TTTAAAAGTG CTCATCATTG GAAAACGTTC TTCGGGGCGA AAACTCTCAA GGATCTTACC GCTGTTGAGA TCCAGTTCGA TGTAACCCAC 5701 5801 TCGTGCACCC AACTGATCTT CAGCATCTTT TACTTTCACC AGCGTTTCTG GGTGAGCAAA AACAGGAAGG CAAAATGCCG CAAAAAAGGG AATAAGGGCG 5851 ACACGGAAAT GTTGAATACT CATACTCTTC CTTTTTCAAT ATTATTGAAG CATTTATCAG GGTTATTGTC TCATGAGCGG ATACATATTT GAATGTATTT 5951 AGAAAAATAA ACAAATAGGG GTTCCGCGCA CATTTCCCCG AAAAGTGCCA 6001 CCTGACGTCT AAGAAACCAT TATTATCATG ACATTAACCT ATAAAAATAG 6051 6101 GCGTATCACG AGGCCCTTTC GTCTCGCGCG TTTCGGTGAT GACGGTGAAA ACCTCTGACA CATGCAGCTC CCGGAGACGG TCACAGCTTG TCTGTAAGCG GATGCCGGGA GCAGACAAGC CCGTCAGGGC GCGTCAGCGG GTGTTGGCGG 6151 6201 6251 GTGTCGGGGC TGGCTTAACT ATGCGGCATC AGAGCAGATT GTACTGAGAG TGCACCATAT GCGGTGTGAA ATACCGCACA GATGCGTAAG GAGAAAATAC CGCATCAGGC GCCATTCGCC ATTCAGGCTG CGCAACTGTT GGGAAGGGCG 6351 ATCGGTGCGG GCCTCTTCGC TATTACGCCA GCTGGCGAAA GGGGGATGTG CTGCAAGGCG ATTAAGTTGG GTAACGCCAG GGTTTTCCCA GTCACGACGT 6401 6451 6501 TGTAAAACGA CGGCCAGTGA ATTTCGACCT GCAGTCGACA GAAGCCTTAC GTGACAGCTG GCGAAGAACC ATGGCCAGCT GGTGACAAGC CAAAACAGCT 6551

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6601	CTGGCTCGCA	AAACATGTTC	CCTTGGCTGC	TTTCCACTTC	CCCTTGTGCT
6651	TTGTTTACTT	GTGTCAGCTG	GTTGGCTCCC	TAGGTATGAG	CTCATGCTTG
6701	GCTGGCAGCC	ATCCAGTTTT	AGCCAGCTCT	GCTTTGTTTA	CTTGTGTCAG
6751	CTGGTTGGCT	CCCTAGGTAT	GAGCTCATGC	TTGGCTGGCA	GCCATCCAGT
6801	TTTAGCCAGC	TCCTCCCTAC	CTTCCCTTTT	TTTTATATAT	ACAGGAGGCC
6851	GAGGCCGCCT	CCGCCTCCAA	GCTTACTCAG	AAGTAGTAAG	GGCGTGGAGG
6901	CTTTTTAGGA	GGCCAGGGAA	ATTCCCTTGT	TTTTCCCTTT	TTTGCAGTAA
6951	TTTTTTGCTG	CAAAAAGCTA	A		

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### pD12JCVPshort-hCNTF

Length: 7558

1 GCTAGCGATT TAGGTGACAC TATAGAATCt cgacnnGTCA CCCCTAGAGT CGAGCTGTGA CGGTCCTTAC AATGAAATGC ANCTGGGTTA TCTTCTTCCT GATGGCAGGG GTTACAGGTA AGGGGCTCCC AAGTCCCAAA CTTGAGGGTC 101 CATAAACTCT GTGACAGTGG CAATCACTTT GCCTTTCTTT CTACAGGGGT 151 GAATTCGGCT TTCACAGAGC ATTCACCGCT GACCCCTCAC CGTCGGGACC 201 TCTGTAGCCG CTCTATCTGG CTAGCAAGGA AGATTCGTTC AGACCTTGAC TGCTCTTACG GAATCCTATG TAAGTTGCCT ATTTTGCTGT TATCTGTTTT CCCTTCATCT TTTTTGATCC AGCAACTTAC CATCACGCAT CAGCTCCATT ACCAATTGTG AAAGCTCTAA TCATATAGTC ATTCATATAG GTTATTTGAC 401 ATGGGCCCTT CCCTTGAGGA AACCCATGTG ACTTTATTTT CTTCCTCTGG GCTGTTTAGG AGATGAAGTT ACTTGAATGA GAAAATATAT ATGGAGTTCT 501 AGAAAGGATT GGTTTATATG TCTTGGAGGC TATTTCAAAA TTTATTTGGC 551 CATATATTCT GAATACTACC TAGAACAGAT TAGCCATGGG CCCTNTGGGT 601 651 TNTTCATAAG CCATTGTTCT GAANTTTTTT AGCTTTGTAA ATGAAAGGTT TATGGGATAG GAAGAGTNCT ATGAACGTGG GAGGAATTTG TAAATCCTAC 701 CAATTINING TATATAGCAT TAGCCCCCAC CITTIANTAT TCTGCATCAA AAGTAAGATT GTGTCTAAAG AGAAAGGTNA GCTATCAAAA GGACTCCTAT 851 AANATTCNTT GGAAACTTNT GGAANTGTCA AATTTNTTTG AGCTAATTNT 901 TGGAGTTCCA AANTTTGTCT TNTNACAGTN AAGGGGGANC CCCATTCANA TTTNCCCCCC TNNNGANAAT GCTTGGGGGA AAAAACCTNC CAACCCCNTT GTGGGANGAA GTTTTTTAA NNTTTTAAGG CTNGNNGAAA CNGGNTTTTA ATTTTTGGG NCNANCGCCT NTCCCCGGTA CCAGGAAAAT CAGGACCTNT TTTTGGGGNN GNGCNCCNAC NGGGGGGNAA AANGGGAAAT TTCNTCANAA 1101 1151 AAAATCTTTT CCGnnnnnng tgaagcatca gggcctgaac aagaacatca acctggactc tgcggatggg atgccagtgg caagcactga tcagtggagt 1201 1251 gagetgaceg aggeagageg actecaagag aacetteaag ettategtae

1301 cttccatgtt ttgttggcca ggctcttaga agaccagcag gtgcatttta 1351 ccccaaccga aggtgacttc catcaagcta tacataccct tcttctccaa 1401 gtegetgeet ttgcatacca gatagaggag ttaatgatac teetggaata 1451 caagateece egeaatgagg etgatgggat geetattaat gttggagatg gtggtetett tgagaagaag etgtggggee taaaggtget geaggagett 1501 tcacagtgga cagtaaggtc catccatgac cttcgtttca tttcttctca 1551 1601 tcagactggg atcccagcac gtgggagcca ttatattgct aacaacaaga aaatgtagnn nnngcggccT GCGCCGTCTT TCCCGACGTT AAAGGGATGA AACCACAAGA CTTACCTTCG CTCGGAAGTA AAACGACAAA CACACACAGT 1701 TTTGCCCGTT TTCATGAGAA ATGGGACGTC TGCGCACGAA ACGCGCCGTC 1751 GCTTGAGGAG GACTTGTACA AACACGATCT ATGCAGGTTT CCCCAACTGA 1801 CACAAACCGT GCAACTTGAA ACTCCGCCTG GTCTTCCAG GTCTAGAGGG 1851 1901 GTAACATTTT GTACTGTGTT TGACTCCACG CTCGATCCAC TAGCGAGTGT TAGTAGCGGT ACTGCTGTCT CGTAGCGGAG CATGTTGGCC GTGGGAACAC 1951 CTCCTTGGTA ACAAGGACCC ACGGGGCCGA AAGCCATGTC CTAACGGACC 2001 2051 CAACATGTGT GCAACCCCAG CACGGCAGCT TTACTGTGAA ACCCACTTCA 2101 AGGTGACATT GATACTGGTA CTCAAACACT GGTGACAGGC TAAGGATGCC CTTCAGGTAC CCCGAGGTAA CAAGCGACAC TCGGGATCTG AGAAGGGGAC 2151 TGGGACTTCT TTAAAGTGCC CAGTTTAAAA AGCTTCTACG CCTGAATAGG 2201 2251 TGACCGGAGG CCGGCACCTT TCCTTTTATA ACCACTGAAC ACATGGAAGA 2301 CGCCAAAAAC ATAAAGAAAG GCCCGGCGCC ATTCTATCCT CTAGAGGATG GAACCGCTGG AGAGCAACTG CATAAGGCTA TGAAGAGATA CGCCCTGGTT 2351 CCTGGAACAA TTGCTTTTAC AGATGCACAT ATCGAGGTGA ACATCACGTA 2451 CGCGGAATAC TTCGAAATGT CCGTTCGGTT GGCAGAAGCT ATGAAACGAT 2501 ATGGGCTGAA TACAAATCAC AGAATCGTCG TATGCAGTGA AAACTCTCTT 2551 CAATTCTTTA TGCCGGTGTT GGGCGCGTTA TTTATCGGAG TTGCAGTTGC 2601 GCCCGCGAAC GACATTTATA ATGAACGTGA ATTGCTCAAC AGTATGAACA 2651 TTTCGCAGCC TACCGTAGTG TTTGTTTCCA AAAAGGGGTT GCAAAAAATT

(4)

2701 TTGAACGTGC AAAAAAATT ACCAATAATC CAGAAAATTA TTATCATGGA 2751 TTCTAAAACG GATTACCAGG GATTTCAGTC GATGTACACG TTCGTCACAT 2801 CTCATCTACC TCCCGGTTTT AATGAATACG ATTTTGTACC AGAGTCCTTT 2851 GATCGTGACA AAACAATTGC ACTGATAATG AATTCCTCTG GATCTACTGG 2901 GTTACCTAAG GGTGTGGCCC TTCCGCATAG AACTGCCTGC GTCAGATTCT 2951 CGCATGCCAG AGATCCTATT TTTGGCAATC AAATCATTCC GGATACTGCG 3001 ATTTTAAGTG TTGTTCCATT CCATCACGGT TTTGGAATGT TTACTACACT 3051 CGGATATTTG ATATGTGGAT TTCGAGTCGT CTTAATGTAT AGATTTGAAG 3101 AAGAGCTGTT TTTACGATCC CTTCAGGATT ACAAAATTCA AAGTGCGTTG 3151 CTAGTACCAA CCCTATTTC ATTCTTCGCC AAAAGCACTC TGATTGACAA 3201 ATACGATTTA TCTAATTTAC ACGAAATTGC TTCTGGGGGC GCACCTCTTT 3251 CGAAGAAGT CGGGGAAGCG GTTGCAAAAC GCTTCCATCT TCCAGGGATA 3301 CGACAGGAT ATGGGCTCAC TGAGACTACA TCAGCTATTC TGATTACACC 3351 CGAGGGGGAT GATAAACCGG GCGCGGTCGG TAAAGTTGTT CCATTTTTTG 3401 AAGCGAAGGT TGTGGATCTG GATACCGGGA AAACGCTGGG CGTTAATCAG 3451 AGAGGCGAAT TATGTGTCAG AGGACCTATG ATTATGTCCG GTTATGTAAA 3501 CAATCCGGAA GCGACCAACG CCTTGATTGA CAAGGATGGA TGGCTACATT 3551 CTGGAGACAT AGCTTACTGG GACGAAGACG AACACTTCTT CATAGTTGAC 3601 CGCTTGAAGT CTTTAATTAA ATACAAAGGA TATCAGGTGG CCCCCGCTGA 3651 ATTGGAATCG ATATTGTTAC AACACCCCAA CATCTTCGAC GCGGGCGTGG 3701 CAGGTCTTCC CGACGATGAC GCCGGTGAAC TTCCCGCCGC CGTTGTTGTT 3751 TTGGAGCACG GAAAGACGAT GACGGAAAAA GAGATCGTGG ATTACGTCGC 3801 CAGTCAAGTA ACAACCGCGA AAAAGTTGCG CGGAGGAGTT GTGTTTGTGG 3851 ACGAAGTACC GAAAGGTCTT ACCGGAAAAC TCGACGCAAG AAAAATCAGA 3901 GAGATCCTCA TAAAGGCCAA GAAGGGCGGA AAGTCCAAAT TGTAAAATGT 3951 AACTGTATTC AGCGATGACG AAATTCTTAG CTATTGTAAT GACTCTAGAG 4001 GATCTTTGTG AAGGAACCTT ACTTCTGTGG TGTGACATAA TTGGACAAAC 4051 TACCTACAGA GATTTAAAGC TCTAAGGTAA ATATAAAATT TTTAAGTGTA

 $(\widehat{\mathbb{S}})$ 

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TAATGTGTTA AACTACTGAT TCTAATTGTT TGTGTATTTT AGATTCCAAC 4151 CTATGGAACT GATGAATGGG AGCAGTGGTG GAATGCCTTT AATGAGGAAA 4201 ACCTGTTTTG CTCAGAAGAA ATGCCATCTA GTGATGATGA GGCTACTGCT 4251 GACTCTCAAC ATTCTACTCC TCCAAAAAAG AAGAGAAAGG TAGAAGACCC 4301 CAAGGACTTT CCTTCAGAAT TGCTAAGTTT TTTGAGTCAT GCTGTGTTTA 4351 GTAATAGAAC TCTTGCTTGC TTTGCTATTT ACACCACAAA GGAAAAAGCT 4401 GCACTGCTAT ACAAGAAAAT TATGGAAAAA TATTCTGTAA CCTTTATAAG 4451 TAGGCATAAC AGTTATAATC ATAACATACT GTTTTTTCTT ACTCCACACA 4501 GGCATAGAGT GTCTGCTATT AATAACTATG CTCAAAAATT GTGTACCTTT 4551 AGCTTTTAA TTTGTAAAGG GGTTAATAAG GAATATTTGA TGTATAGTGC 4601 CTTGACTAGA GATCATAATC AGCCATACCA CATTTGTAGA GGTTTTACTT 4651 GCTTTAAAAA ACCTCCCACA CCTCCCCCTG AACCTGAAAC ATAAAATGAA 4701 TGCAATTGTT GTTGTTAACT TGTTTATTGC AGCTTATAAT GGTTACAAAT 4751 AAAGCAATAG CATCACAAAT TTCACAAATA AAGCATTTTT TTCACTGCAT 4801 TCTAGTTGTG GTTTGTCCAA ACTCATCAAT GTATCTTATC ATGTCTGGAT 4851 CCCCGGGTCC CTATAGTGAG TCGTATTAGC TTGGCGTAAT CATGGTCATA 4901 GCTGTTTCCT GTGTGAAATT GTTATCCGCT CACAATTCCA CACAACATAC 4951 GAGCCGGAAG CATAAAGTGT AAAGCCTGGG GTGCCTAATG AGTGAGCTAA 5001 CTCACATTAA TTGCGTTGCG CTCACTGCCC GCTTTCCAGT CGGGAAACCT 5051 GTCGTGCCAG CTGCATTAAT GAATCGGCCA ACGCGCGGG AGAGGCGGTT 5101 TGCGTATTGG GCGCTCTTCC GCTTCCTCGC TCACTGACTC GCTGCGCTCG 5151 GTCGTTCGGC TGCGGCGAGC GGTATCAGCT CACTCAAAGG CGGTAATACG 5201 GTTATCCACA GAATCAGGGG ATAACGCAGG AAAGAACATG TGAGCAAAAG 5251 GCCAGCAAAA GGCCAGGAAC CGTAAAAAGG CCGCGTTGCT GGCGTTTTTC 5301 CATAGGCTCC GCCCCCTGA CGAGCATCAC AAAAATCGAC GCTCAAGTCA 5351 GAGGTGGCGA AACCCGACAG GACTATAAAG ATACCAGGCG TTTCCCCCTG 5401 GAAGCTCCCT CGTGCGCTCT CCTGTTCCGA CCCTGCCGCT TACCGGATAC 5451 CTGTCCGCCT TTCTCCCTTC GGGAAGCGTG GCGCTTTCTC AATGCTCACG

5501 CTGTAGGTAT CTCAGTTCGG TGTAGGTCGT TCGCTCCAAG CTGGGCTGTG 5551 TGCACGAACC CCCCGTTCAG CCCGACCGCT GCGCCTTATC CGGTAACTAT CGTCTTGAGT CCAACCCGGT AAGACACGAC TTATCGCCAC TGGCAGCAGC 5601 5651 CACTGGTAAC AGGATTAGCA GAGCGAGGTA TGTAGGCGGT GCTACAGAGT TCTTGAAGTG GTGGCCTAAC TACGGCTACA CTAGAAGGAC AGTATTTGGT 5701 ATCTGCGCTC TGCTGAAGCC AGTTACCTTC GGAAAAAGAG TTGGTAGCTC TTGATCCGGC AAACAAACCA CCGCTGGTAG CGGTGGTTTT TTTGTTTGCA AGCAGCAGAT TACGCGCAGA AAAAAAGGAT CTCAAGAAGA TCCTTTGATC 5901 TTTTCTACGG GGTCTGACGC TCAGTGGAAC GAAAACTCAC GTTAAGGGAT 5951 TTTGGTCATG AGATTATCAA AAAGGATCTT CACCTAGATC CTTTTAAATT 6001 AAAAATGAAG TTTTAAATCA ATCTAAAGTA TATATGAGTA AACTTGGTCT 6051 GACAGTTACC AATGCTTAAT CAGTGAGGCA CCTATCTCAG CGATCTGTCT 6101 ATTTCGTTCA TCCATAGTTG CCTGACTCCC CGTCGTGTAG ATAACTACGA TACGGGAGGG CTTACCATCT GGCCCCAGTG CTGCAATGAT ACCGCGAGAC 6151 6201 CCACGCTCAC CGGCTCCAGA TTTATCAGCA ATAAACCAGC CAGCCGGAAG 6251 GGCCGAGCGC AGAAGTGGTC CTGCAACTTT ATCCGCCTCC ATCCAGTCTA 6301 TTAATTGTTG CCGGGAAGCT AGAGTAAGTA GTTCGCCAGT TAATAGTTTG CGCAACGTTG TTGCCATTGC TACAGGCATC GTGGTGTCAC GCTCGTCGTT TGGTATGGCT TCATTCAGCT CCGGTTCCCA ACGATCAAGG CGAGTTACAT GATCCCCCAT GTTGTGCAAA AAAGCGGTTA GCTCCTTCGG TCCTCCGATC 6501 GTTGTCAGAA GTAAGTTGGC CGCAGTGTTA TCACTCATGG TTATGGCAGC ACTGCATAAT TCTCTTACTG TCATGCCATC CGTAAGATGC TTTTCTGTGA 6551 CTGGTGAGTA CTCAACCAAG TCATTCTGAG AATAGTGTAT GCGGCGACCG AGTTGCTCTT GCCCGGCGTC AATACGGGAT AATACCGCGC CACATAGCAG 6651 6701 AACTTTAAAA GTGCTCATCA TTGGAAAACG TTCTTCGGGG CGAAAACTCT 6751 CAAGGATCTT ACCGCTGTTG AGATCCAGTT CGATGTAACC CACTCGTGCA 6801 CCCAACTGAT CTTCAGCATC TTTTACTTTC ACCAGCGTTT CTGGGTGAGC 6851 AAAAACAGGA AGGCAAAAATG CCGCAAAAAA GGGAATAAGG GCGACACGGA

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AATGTTGAAT ACTCATACTC TTCCTTTTC AATATTATTG AAGCATTTAT

CAGGGTTATT GTCTCATGAG CGGATACATA TTTGAATGTA TTTAGAAAAA

CAGAGGTTATT GTCTCATGAG CGGATACATA TTTGAATGTA TTTAGAAAAA

TOO1 TAAACAAATA GGGGTTCCGC GCACATTTCC CCGAAAAGTG CCACCTGACG

TCTAAAGAAAC CATTATTATC ATGACATTAA CCTATAAAAA TAGGCGTATC

ACGAGGCCCT TTCGTCTCGC GCGTTTCGGT GATGACGGTG AAAACCTCTG

ACACATGCAG CTCCCGGAGA CGGTCACAGC TTGTCTGTAA GCGGATGCCG

GGAGCAGACA AGCCCGTCAG GGCGCGTCAG CGGGTGTTGG CGGGTGTCGG

GGCTGGCTTA ACTATGCGGC ATCAGAGCAG ATTGTACTGA GAGTGCACCA

TATGCGGTGT GAAATACCGC ACAGATGCGT AAGGAGAAAA TACCGCATCA

TATGCGGTGT GAAATACCGC CTGCGCAACT GTTGGGAAGG GCGATCGGTG

CGGCCCATTC GCCATTCAGG CTGCGCAACT GTTGGGAAGG GCGATCGGTG

CGGCCCTCTT CGCTATTACG CCAGCTGGCG AAAGGGGGAT GTGCTGCAAG

CGACGGCCAG TGAATTTCGA CCTGCAGLCG actttttta tatatacagg

TSS1 aggccgag

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JCVPshort-hgdnf Length: 6565 June 8, 1999 16:57 Type: N Check:

	1 51	GCTAGCGATT	TAGGTGACAC	TATAGAATAG	G ATCCCCATGA	AGTTATGGGA
		中で中でで中ででです				
		IGICGIGGCI	' GTCTGCCTGG	TGCTGCTCC	A CACCGCGTCC	GCCTTCCCGC
	101	TGCCCGCCGG	F TAAGAGGCCT	CCCGAGGCGC	CCGCCGAAGA	CCGCTCCCTC
	151	GGCCGCCGCC	GCGCGCCCTT	CGCGCTGAG	2 AGTGACTCAA	ATATGCCAGA
	201	GGATTATCCT	GATCAGTTCC	ATGATGTCAT	r ggattttatt	CAAGCCACCA
	251	TTAAAAGACI	' GAAAAGGTCA	CCAGATAAA	2 AAATGGCAGT	GCTTCCTAGA
	301	AGAGAGCGGA	ATCGGCAGGC	TGCAGCTGC	AACCCAGAGA	ATTCCAGAGG
	351	AAAAGGTCGG	AGAGGCCAGA	GGGGCAAAA	CCGGGGTTGT	GTCTTAACTG
	401	CAATACATTT	AAATGTCACT	GACTTGGGTC	TGGGCTATGA	AACCAAGGAG
	451	GAACTGATTT	TTAGGTACTG	CAGCGGCTCI	TGCGATGCAG	CTGAGACAAC
	501 551	GTACGACAAA	ATATTGAAAA	ACTTATCCAG	AAATAGAAGG	CTGGTGAGTG
	601	TCCTTTTTT	GCAGGCATGT	TGCAGACCCA	TCGCCTTTGA	TGATGACCTG
	651	TOGITITIAG	CCATCHARCCT	GGTTTACCAT	ATTCTAAGAA	AGCATTCCGC
	701	GGGATGAAAC	CACAACACTE	ACCURCOCO	CCGTCTTTCC GGAAGTAAAA	CGACGTTAAA
	751	ACACACTTTT	CACAAGACII	ACCITCGCTC	GGACGTCTGC	CGACAAACAC
	801	CGCCGTCGCT	TGAGGAGGAG	TTCTACAAAIG	ACGATCTATG	GCACGAAACG
	851	CAACTGACAC	AAACCGTGCA	ACTTCAAAC	CCGCCTGGTC	TORROGATOR
	901	TAGAGGGGTA	ACATTTTGTA	CTCTCTTTCA	CTCCACGCTC	CATCCAGGIC
	951	CGAGTGTTAG	TAGCGGTACT	GCTGTCTCGT	AGCGGAGCAT	CTTCCCCCTC
1	001	GGAACACCTC	CTTGGTAACA	AGGACCCACG	GGGCCGAAAG	CCATCTCCTA
1	051	ACGGACCCAA	CATGTGTGCA	ACCCCAGCAC	GGCAGCTTTA	CTGTGAAACC
1	101	CACTTCAAGG	TGACATTGAT	ACTGGTACTC	AAACACTGGT	GACAGGCTAA
1	151	GGATGCCCTT	CAGGTACCCC	GAGGTAACAA	GCGACACTCG	GGATCTGAGA
	201	AGGGGACTGG	GACTTCTTTA	AAGTGCCCAG	TTTAAAAAGC	TTCTACGCCT
	251	GAATAGGTGA	CCGGAGGCCG	GCACCTTTCC	TTTTATAACC	ACTGAACACA
	301	TGGAAGACGC	CAAAAACATA	AAGAAAGGCC	CGGCGCCATT	CTATCCTCTA
	351	GAGGATGGAA	CCGCTGGAGA	GCAACTGCAT	AAGGCTATGA	AGAGATACGC
	401	CCTGGTTCCT	GGAACAATTG	CTTTTACAGA	TGCACATATC	GAGGTGAACA
	451	TCACGTACGC	GGAATACTTC	GAAATGTCCG	TTCGGTTGGC	AGAAGCTATG
	501 551	AAACGATATG	GGCTGAATAC	AAATCACAGA	ATCGTCGTAT	GCAGTGAAAA
	601	CACTUTUTAA	TTCTTTATGC	CGGTGTTGGG	CGCGTTATTT	ATCGGAGTTG
	651	ATCAACATT	CGCGAACGAC	ATTTATAATG	AACGTGAATT	GCTCAACAGT
	701	AIGMACAIII	AACCTCCAAA	CGTAGTGTTT	GTTTCCAAAA AATAATCCAG	AGGGGTTGCA
	751	TCATCCATTC	TAAAAACCAAA	TACCACCCAT	TTCAGTCGAT	AAAATTATTA
	801	GTCACATCTC	ATCTACCTCC	CCCTTTTANT	GAATACGATT	GTACACGTTC
	851	GTCCTTTGAT	CGTGACAAAA	CAATTGCACT	GATAATGAAT	TIGIACCAGA
	901	CTACTGGGTT	ACCTAAGGGT	GTGGCCCTTC	CGCATAGAAC	TCCTCTGGAT
19	951	AGATTCTCGC	ATGCCAGAGA	TCCTATTTT	GGCAATCAAA	TCATTCCCCA
20	001	TACTGCGATT	TTAAGTGTTG	TTCCATTCCA	TCACGGTTTT	GGAATGTTTA
	051	CTACACTCGG	ATATTTGATA	TGTGGATTTC	GAGTCGTCTT	AATGTATAGA
21	101	TTTGAAGAAG	AGCTGTTTTT	ACGATCCCTT	CAGGATTACA	AAATTCAAAG
	151	TGCGTTGCTA	GTACCAACCC	TATTTTCATT	CTTCGCCAAA	AGCACTCTGA
	201	TTGACAAATA	CGATTTATCT	AATTTACACG	AAATTGCTTC	TGGGGGGCGCA
	251	CCTCTTTCGA	AAGAAGTCGG	GGAAGCGGTT	GCAAAACGCT	TCCATCTTCC
	301	AGGGATACGA	CAAGGATATG	GGCTCACTGA	GACTACATCA	GCTATTCTGA
	351 101	TTACACCCGA	GGGGGATGAT	AAACCGGGCG	CGGTCGGTAA	AGTTGTTCCA
	151	TITITIGAAG	CGAAGGTTGT	GGATCTGGAT	ACCGGGAAAA	CGCTGGGCGT
	01	ATCTAAACAA	TCCCCAATTAT	GTGTCAGAGG	ACCTATGATT	ATGTCCGGTT
	551	CTACATTCTC	CACACAMACC	TTACTCCCAC	TGATTGACAA GAAGACGAAC	GGATGGATGG
	501	AGTTGACCGC	サヤになるとれてみない	1174C1GGGAC	CAAAGACGAAC A	ACTICTICAT
	551	CCGCTGAATT	GGAATCGATA	TAMIIAMAIA	ACCCCAACAT	CAGGTGGCCC
	01	GGCGTGGCAG	GTCTTCCCGA	CGATGACGCC	GGTGAACTTC (	CCCCCCCCC
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	.01	AAGTGTATAA	TGTGTTAAAC	TACTGATTCT	AATTGTTTGT (	GTATTTTAGA
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**-** 37/56 -

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pRetroOFF-E6E7 Length: 7840 June 10, 1999 12:21 Type: N Check: 5234

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3151
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(3)

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pRetroOFF-U19tsa58 Length: 8852

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puhd10-3-hIL3 Length: 3621

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3551	atactettee	tttttcaata	ttattgaagc	atttatcagg	gttattgtct
3601	catgagcgga	tacatatttg	aatgtattta	gaaaaataaa	caaatagggg
3651	ttccgcgcac	atttccccga	aaagtgccac	ctgacgtcta	agaaaccatt
3701	attatcatga	cattaaccta	taaaaatagg	cgtatcacga	ggccctttcg
3751	t.c				

**(**)

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#### puhd10-3-tgf

(5°)

1	ctcgagtttaccactccctatcagtgatagagaaaagtgaaagtcgagtttaccactccc	60
61	tatcagtgatagagaaaagtgaaagtcgagtttaccactccctatcagtgatagagaaaa	120
121	gtgaaagtcgagtttaccactccctatcagtgatagagaaaagtgaaagtcgagtttacc	180
181	actccctatcagtgatagagaaaagtgaaagtcgagtttaccactccctatcagtgatag	240
241	agaaaagtgaaagtcgagtttaccactccctatcagtgatagagaaaagtgaaagtcgag	300
301	ctcggtacccgggtcgagtaggcgtgtacggtgggaggcctatataagcagagctcgttt	360
361	agtgaaccgtcagatcgcctggagacgccatccacgctgttttgacctccatagaagaca	420
421	ccgggaccgatccagcctccgcggccccgaattcctgcagcccATGCACTTGCAAAGGGC	480
481	TCTGGTAGTCCTGGCCCTGCTGAACTTGGCCACAATCAGCCTCTCTCT	540
541	CACGTTGGACTTCGGCCACATCAAGAAGAAGAGGGTGGAAGCCATTAGGGGACAGATCTT	600
601	GAGCAAGCTCAGGCTCACCAGCCCCCTGAGCCATCGGTGATGACCCACGTCCCCTATCA	660
661	GGTCCTGGCACTTTACAACAGCACCCGGGAGTTGCTGGAAGAGATGCACGGGGAGAGGGA	720
721	GGAAGGCTGCACTCAGGAGACCTCGGAGTCTGAGTACTATGCCAAAGAGATCCATAAATT	780
781	CGACATGATCCAGGGACTGGCGGAGCACAATGAACTGGCCGTCTGCCCCAAAGGAATTAC	840
841	CTCTAAGGTTTTTCGTTTCAATGTGTCCTCAGTGGAGAAAAATGGAACCAATCTGTTCCG	900
901	GGCAGAGTTCCGGGTCTTGCGGGTGCCCAACCCCAGCTCCAAGCGCACAGAGCAGAGAAT	960
961	TGAGCTCTTCCAGATACTTCGACCGGATGAGCACATAGCCAAGCAGCGCTACATAGGTGG	1020
1021	CAAGAATCTGCCCACAAGGGGCACCGCTGAATGGCTGTCTTTCGATGTCACTGACACTGT	1080
1081	GCGCGAGTGGCTGTTGAGGAGAGAGTCCAACTTGGGTCTGGAAATCAGCATCCACTGTCC	1140
1141	ATGTCACACCTTTCAGCCCAATGGAGACATACTGGAAAATGTTCATGAGGTGATGGAAAT	1200
1201	CAAATTCAAAGGAGTGGACAATGAAGATGACCATGGCCGTGGAGACCTGGGGCGTCTCAA	1260
1261	GAAGCAAAAGGATCACCACAACCCACACCTGATCCTCATGATGATCCCCCCACACCGACT	1320

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1321	GGACAGCCCAGGCCAGGCAGTCAGAGGAGAAGAAGAGGCCCTGGACACCAATTACTGCTT	1380
1381	CCGCAACCTGGAGGAGAACTGCTGTGTACGCCCCCTTTATATTGACTTCCGGCAGGATCT	1440
1441	AGGCTGGAAATGGGTCCACGAACCTAAGGGTTACTATGCCAACTTCTGCTCAGGCCCTTG	1500
1501	CCCATACCTCCGCAGCGCAGACACCACATAGCACGGTGCTTGGACTATACAACACCCT	1560
1561	GAACCCAGAGCGTCTGCCTCGCCATGCTGCGTCCCCCAGGACCTGGAGCCCCTGACCAT	1620
1621	CTTGTACTATGTGGGCAGAACCCCCAAGGTGGAGCAGCTGTCCAACATGGTGGTGAAGTC	1680
1681	GTGTAAGTGCAGCTGAgggggatccactagttctagaggatccagacatgataagataca	1740
1741	ttgatgagtttggacaaaccacaactagaatgcagtgaaaaaaatgctttatttgtgaaa	1800
1801	tttgtgatgctattgctttatttgtaaccattataagctgcaataaacaagttaacaaca	1860
1861	acaattgcattcattttatgtttcaggttcaggggggggtgtggggaggttttttaaagca	1920
1921	agtaaaacctctacaaatgtggtatggctgattatgatcctgcaagcctcgtctggc	1980
L981	cggaccacgctatctgtgcaaggtccccggacgcgctccatgagcagagcgcccgcc	2040
2041	ccgaggcaagactcgggcggccctgccgtcccaccaggtcaacaggcggtaaccggc	2100
2101	ctcttcatcgggaatgcgcgcgaccttcagcatcgccggcatgtcccctggcggacggga	2160
2161	agtatcagctcgaccaagcttggcgagattttcaggagctaaaggaagctaaaatggagaa	2220
2221	aaaaatcactggatataccaccgttgatatatcccaatggcatcgtaaagaacattttga	2280
2281	ggcatttcagtcagttgctcaatgtacctataaccagaccgttcagctgcattaatgaat	2340
2341	cggccaacgcgcggggagaggcggtttgcgtattgggcgctcttccgcttcctcgctcac	2400
2401	tgactcgctgcgctcggtcgttcggctgcggcggagcggtatcagctcactca	2460
	aatacggttatccacagaatcaggggataacgcaggaaagaacatgtgagcaaaaggcca	2520
2521		2580
2581	ccctgacgagcatcacaaaaatcgacgctcaagtcagaggtggcgaaacccgacaggact	2640

## - 50/56 -

264	ataaagataccaggcgtttccccctggaagctccctcgtgcgctctcctgttccgaccc 1	t + 2700
270	gccgcttaccggatacctgtccgcctttctcccttcgggaagcgtggcgctttctcaatc	g + 2760
276	ctcacgctgtaggtatctcagttcggtgtaggtcgttcgctccaagctgggctgtgtgc	a + 2820
282	cgaaccccccgttcagcccgaccgctgcgccttatccggtaactatcgtcttgagtccaa	a + 2880
2881	cccggtaagacacgacttatcgccactggaagcagccactggtaacaggattagcagagc	: - 2940
2941	gaggtatgtaggcggtgctacagagttcttgaagtggtggcctaactacggctacactac 	f - 3000
3001	aaggacagtatttggtatctgcgctctgctgaagccagttaccttcggaaaaagagttgg	3060
	tagetettgateeggeaaacaaaceacegetggtageggtggttttttttgtttgcaagea	
	gcagattacgcgcagaaaaaaggatctcaagaagatcctttgatcttttctacggggtc	
	tgacgctcagtggaacgaaaactcacgttaagggattttggtcatgagattatcaaaaag	
	gatetteacetagateettttaaattaaaaatgaagttttaaateaate	
	tgagtaaacttggtctgacagttaccaatgcttaatcagtgaggcacctatctcagcgat	
	ctgtctatttcgttcatccatagttgcctgactccccgtcgtgtagataactacgatacg	
3421	ggagggcttaccatetggccccagtgctgcaatgataccgcgagacccacgctcaccggc	3420
	tccagatttatcagcaataaaccagccagccggaagggccgagcgcagaagtggtcctgc	
	aactttatccgcctccatccagtctattaattgttgccgggaaggtagagtagatagttg	
	gccagttaatagtttgcgcaacgttgttgccattgctacaggcatcgtgtggtcacgctc	
	gtcgtttggtatggcttcattcagctccggttcccaacgatcaaggcgagttacatgatc	
	ccccatgttgtgcaaaaaagcggttagctccttcggtcctccgatcgttgtcagaagtaa	
	gttggccgcagtgttatcactcatggttatggcagcactgcataattctcttactgtcat	
	qccatccqtaagatgcttttctqtgactqqtgaqtactcaacqaagtaattatgagaata	
841	qtqtatqcqqcqacqaqttqctcttqccqtcqtcatcatacqqqataatacqqqqa	
901	tagcagaactttaaaagtgctcatcattggaaaacgttcttcggggcgaaaactctcaag	3960

(<sub>650</sub>)

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# - 51/56 -

3961		4020
4021	gatettacegetgttgagatecagttegatgtaacecactegtgeacecaactgatette	4080
4081	agcatcttttactttcaccagcgtttctgggtgagcaaaacaggaaggcaaaatgccgc	4140
4141	aaaaaagggaataagggcgacacggaaatgttgaatactcatactcttcctttttcaata	4200
4201	ttattgaagcatttatcagggttattgtctcatgagcggatacatatttgaatgtattta	4260
4261	gaaaaataaacaaataggggttccgcgcacatttccccgaaaagtgccacctgacgtcta	4320
4321	agaaaccattattatcatgacattaacctataaaaataggcgtatcacgaggccctttcg	4380
	tc 4382	

# pUHD10.3-hflt3 Ligand-exon 6 plasmid Length: 4224

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- 1 CTCGAGTTTA CCACTCCCTA TCAGTGATAG AGAAAAGTGA AAGTCGAGTT
- 51 TACCACTCCC TATCAGTGAT AGAGAAAGT GAAAGTCGAG TTTACCACTC
- 101 CCTATCAGTG ATAGAGAAAA GTGAAAGTCG AGTTTACCAC TCCCTATCAG
- 151 TGATAGAGAA AAGTGAAAGT CGAGTTTACC ACTCCCTATC AGTGATAGAG
- 201 AAAAGTGAAA GTCGAGTTTA CCACTCCCTA TCAGTGATAG AGAAAAGTGA
- 251 AAGTCGAGTT TACCACTCCC TATCAGTGAT AGAGAAAAGT GAAAGTCGAG
- 301 CTCGGTACCC GGGTCGAGTA GGCGTGTACG GTGGGAGGCC TATATAAGCA
- 351 GAGCTCGTTT AGTGAACCGT CAGATCGCCT GGAGACGCCA TCCACGCTGT
- 401 TTTGACCTCC ATAGAAGACA CCGGGACCGA TCCAGCCTCC GCGGCCCCGA
- 451 ATTCCggggc ccccggccga aATGacagtg ctggcgccag cctggagccc
- 501 aacaacetat eteeteetge tgetgetget gageteggga eteagtggga
- 551 cccaggactg etectteeaa eacageecea teteeteega ettegetgte
- 601 aaaatccgtg agetgtetga etacetgett caagattace cagtcaccgt
- 651 ggcctccaac ctgcaggacg aggagctctg cgggggcctc tggcggctgg
- 701 tcctggcaca gcgctggatg gagcggctca agactgtcgc tgggtccaag
- 751 atgcaagget tgetggageg egtgaacaeg gagatacaet ttgtcaccaa
- 801 atgtgccttt cageccccc ccagetgtct tegettegte cagaccaaca
- 851 tetecegeet cetgeaggag aceteegage agetggtgge getgaageee
- 901 tggatcactc gccagaactt ctcccggtgc ctggagctgc agtgtcagcc
- 951 cgtagagacg gtgtttcacc gtgtcagcca ggatggtctc gatctcctga
- 1001 cctcgTGAtc tgcccgcctc ggcctcccaa agtgctagga ttacagatac
- 1051 tecteaacce tgecaccee atggagteec eggeceetgg aggecacage
- 1101 cccgacagec ccgcageccc etetgeteet ectactgetg etgecegtgg
- 1151 geeteetget getggeeget geetggtgee tgeaetggea gaggaegegg
- 1201 cggaggacae eccgecetgg ggagcaggtg ecceegtee ecagteecea
- 1251 ggacctgctg cttgtggagc actgacctgg ccaaggcctc atcctgcgga
- 1301 gccttaaaca acgcagtgag acagacatct atcatcccat tttacagggg
- 1351 aggatactga ggcacacaga ggggagtcac cagccagagg atgtatagcc
- 1401 tggacacaga ggaagttggc tagaggccgg tcccttcctt gggcccctct
- 1451 cattecetee ceagaatgga ggeaacgeea gaateeagea eeggeeecat
- 1501 ttacccaact etgaacaaag cccCCGGAAT TCGAGCTCGG TACCCGGGGA

'b: ...

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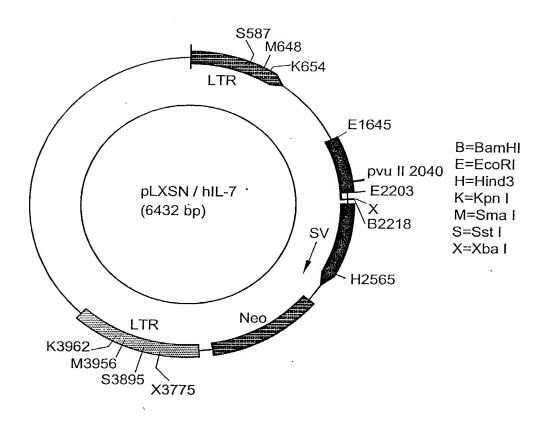
1551 TCCTCTAGAG GATCCAGACA TGATAAGATA CATTGATGAG TTTGGACAAA 1601 CCACAACTAG AATGCAGTGA AAAAAATGCT TTATTTGTGA AATTTGTGAT 1651 GCTATTGCTT TATTTGTAAC CATTATAAGC TGCAATAAAC AAGTTAACAA 1701 CAACAATTGC ATTCATTTA TGTTTCAGGT TCAGGGGGAG GTGTGGGAGG 1751 TTTTTTAAAG CAAGTAAAAC CTCTACAAAT GTGGTATGGC TGATTATGAT 1801 CCTGCAAGCC TCGTCGTCTG GCCGGACCAC GCTATCTGTG CAAGGTCCCC 1851 GGACGCGCG TCCATGAGCA GAGCGCCCGC CGCCGAGGCA AGACTCGGGC 1901 GGCGCCTGC CCGTCCCACC AGGTCAACAG GCGGTAACCG GCCTCTTCAT 1951 CGGGAATGCG CGCGACCTTC AGCATCGCCG GCATGTCCCC TGGCGGACGG 2001 GAAGTATCAG CTCGACCAAG CTTGGCGAGA TTTTCAGGAG CTAAGGAAGC 2051 TAAAATGGAG AAAAAAATCA CTGGATATAC CACCGTTGAT ATATCCCAAT 2101 GGCATCGTAA AGAACATTTT GAGGCATTTC AGTCAGTTGC TCAATGTACC 2151 TATAACCAGA CCGTTCAGCT GCATTAATGA ATCGGCCAAC GCGCGGGGAG 2201 AGGCGGTTTG CGTATTGGGC GCTCTTCCGC TTCCTCGCTC ACTGACTCGC 2251 TGCGCTCGGT CGTTCGGCTG CGGCGAGCGG TATCAGCTCA CTCAAAGGCG 2301 GTAATACGGT TATCCACAGA ATCAGGGGAT AACGCAGGAA AGAACATGTG 2351 AGCAAAAGGC CAGCAAAAGG CCAGGAACCG TAAAAAAGGCC GCGTTGCTGG 2401 CGTTTTTCCA TAGGCTCCGC CCCCTGACG AGCATCACAA AAATCGACGC 2451 TCAAGTCAGA GGTGGCGAAA CCCGACAGGA CTATAAAGAT ACCAGGCGTT 2501 TCCCCTGGA AGCTCCCTCG TGCGCTCTCC TGTTCCGACC CTGCCGCTTA 2551 CCGGATACCT GTCCGCCTTT CTCCCTTCGG GAAGCGTGGC GCTTTCTCAA 2601 TGCTCACGCT GTAGGTATCT CAGTTCGGTG TAGGTCGTTC GCTCCAAGCT 2651 GGGCTGTGTG CACGAACCCC CCGTTCAGCC CGACCGCTGC GCCTTATCCG 2701 GTAACTATCG TCTTGAGTCC AACCCGGTAA GACACGACTT ATCGCCACTG 2751 GCAGCAGCCA CTGGTAACAG GATTAGCAGA GCGAGGTATG TAGGCGGTGC 2801 TACAGAGTTC TTGAAGTGGT GGCCTAACTA CGGCTACACT AGAAGGACAG 2851 TATTTGGTAT CTGCGCTCTG CTGAAGCCAG TTACCTTCGG AAAAAGAGTT 2901 GGTAGCTCTT GATCCGGCAA ACAAACCACC GCTGGTAGCG GTGGTTTTTT 2951 TGTTTGCAAG CAGCAGATTA CGCGCAGAAA AAAAGGATCT CAAGAAGATC 3001 CTTTGATCTT TTCTACGGGG TCTGACGCTC AGTGGAACGA AAACTCACGT 3051 TAAGGGATTT TGGTCATGAG ATTATCAAAA AGGATCTTCA CCTAGATCCT 3101 TTTAAATTAA AAATGAAGTT TTAAATCAAT CTAAAGTATA TATGAGTAAA 3151 CTTGGTCTGA CAGTTACCAA TGCTTAATCA GTGAGGCACC TATCTCAGCG

" b! ...

(

3201 ATCTGTCTAT TTCGTTCATC CATAGTTGCC TGACTCCCCG TCGTGTAGAT 3251 AACTACGATA CGGGAGGGCT TACCATCTGG CCCCAGTGCT GCAATGATAC 3301 CGCGAGACCC ACGCTCACCG GCTCCAGATT TATCAGCAAT AAACCAGCCA 3351 GCCGGAAGGG CCGAGCGCAG AAGTGGTCCT GCAACTTTAT CCGCCTCCAT 3401 CCAGTCTATT AATTGTTGCC GGGAAGCTAG AGTAAGTAGT TCGCCAGTTA 3451 ATAGTTTGCG CAACGTTGTT GCCATTGCTA CAGGCATCGT GGTGTCACGC 3501 TCGTCGTTTG GTATGGCTTC ATTCAGCTCC GGTTCCCAAC GATCAAGGCG 3551 AGTTACATGA TCCCCCATGT TGTGCAAAAA AGCGGTTAGC TCCTTCGGTC 3601 CTCCGATCGT TGTCAGAAGT AAGTTGGCCG CAGTGTTATC ACTCATGGTT 3651 ATGGCAGCAC TGCATAATTC TCTTACTGTC ATGCCATCCG TAAGATGCTT 3701 TTCTGTGACT GGTGAGTACT CAACCAAGTC ATTCTGAGAA TAGTGTATGC 3751 GGCGACCGAG TTGCTCTTGC CCGGCGTCAA TACGGGATAA TACCGCGCCA 3801 CATAGCAGAA CTTTAAAAGT GCTCATCATT GGAAAACGTT CTTCGGGGCG 3851 AAAACTCTCA AGGATCTTAC CGCTGTTGAG ATCCAGTTCG ATGTAACCCA 3901 CTCGTGCACC CAACTGATCT TCAGCATCTT TTACTTTCAC CAGCGTTTCT 3951 GGGTGAGCAA AAACAGGAAG GCAAAATGCC GCAAAAAAGG GAATAAGGGC 4001 GACACGGAAA TGTTGAATAC TCATACTCTT CCTTTTTCAA TATTATTGAA 4051 GCATTTATCA GGGTTATTGT CTCATGAGCG GATACATATT TGAATGTATT 4101 TAGAAAAATA AACAAATAGG GGTTCCGCGC ACATTTCCCC GAAAAGTGCC 4151 ACCTGACGTC TAAGAAACCA TTATTATCAT GACATTAACC TATAAAAATA 4201 GGCGTATCAC GAGGCCCTTT CGTC

- 55/56 -Recovery of insert: EcoRI



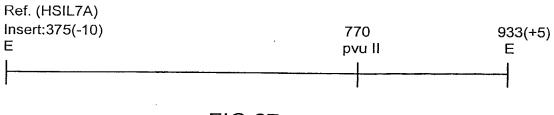


FIG.27

# - 56/56 -

## Plasmid-chart

Designation:

pLXSN/hIL-2

Log no.:

Insert:

( S. )

hll-2 (473bp)

Location:

Vector:

pLXSN (5874bp)

Selection: Amp

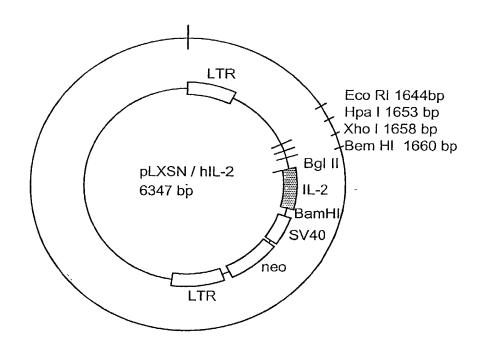
Recovery of insert: Eco RI /Bam HI

Ref.: pLXSN BioTechniques 7,980-987(1989)

Hpal / Bam HI

Xho I / Bam HI

hlL-2 Nature 302,305-309(1983)



Insert: Bgl II 5' AGA TCT ACA - IL-2 - TAA TTA AGT BamHI 473 bp

FIG.28